

Sequence Listing

<110> CHOE, Mu-Hyeon

<120> THE DIMER OF CHIMERIC RECOMBINANT BINDING DOMAIN-FUNCTIONAL GROUP
FUSION FORMED VIA DISULFIDE-BOND-BRIDGE AND THE PROCESSES FOR
PRODUCING THE SAME

<130> YL04011PCT

<140> PCT/KR2004/001595

<141> 2004-06-30

<150> KR2003-0043599

<151> 2003-06-30

<160> 12

<170> KopatentIn 1.71

<210> 1

<211> 1749

<212> DNA

<213> Artificial Sequence

<220>

<223> pMC74 plasmid coding sequence

<400> 1

atgggatgtga agctggtgga atctggagga ggcttagtgc agcctggagg gtccctgaaa 60

ctctcctgtg caacctctgg attcactttc agtgactatt acatgtattg ggttcgccag 120

actccagaga agaggctgga gtgggtcgca tacattagta atgatgatag ttccgccgct 180

tattcagaca ctgtaaaggg ccggttcacc atctccagag acaatgccag gaacaccctc 240

tacctgcaaa tgagccgtct gaagtctgag gacacagcca tatattcctg tgcaagagga 300

ctggcctggg gagcctggtt tgcttactgg ggccaaggga ctctggtcac tgtctctgca 360

Sequence Listing

gccaaaacga ccccccatc tgtctatcca ctggccctg gatctgctgc ccaaactaac	420
tccatggtga ccctgggatg cctggtcaag ggctatttcc ctgagccagt gacagtgacc	480
tggaactctg gatccctgtc cagcgggtgtg cacaccttcc cagctgtcct gcagtctgac	540
ctctacactc tgagcagctc agtgactgtc ccctccagca cctggcccag cgagaccgtc	600
acctgcaacg ttgccacccc ggccagcagc accaagggtgg acaagaaaat tgtgcccagg	660
gattgtggtg gtaagcctag cataagtaca aaagcttccg gaggtccga gggcggcagc	720
ctggccgcgc tgaccgcga ccaggcttgc cacctgccgc tggagacttt caccggtcat	780
cgccagccgc gggctggga acaactggag cagtgcggct atccggtgca gcggtgtg	840
gccctctacc tggcgcgcg gctgtcgtgg aaccaggtcg accaggtgat ccgcaacgcc	900
ctggccagcc ccggcagcgg cggcgacctg ggcaagcga tccgcgagca gccggagcag	960
gcccgtctgg ccctgacct ggccgcgcc gagagcgagc gcttcgtccg gcagggcacc	1020
ggcaacgacg aggccggcgc ggccaacggc ccggcggaca gcggcgacgc cctgctggag	1080
cgcaactatc cactggcgc ggagttctc ggcgacggcg gcgacgtcag cttcagcacc	1140
cgcggcacgc agaactggac ggtggagcgg ctgctccagg cgcaccgcca actggaggag	1200
cgcggtatg tgttcgtcgg ctaccacggc accttcctcg aagcggcgca aagcatcgtc	1260
ttcggcgggg tgcgcgcgcg cagccaggac ctgcacgca tctggcgcgg tttctatata	1320
gccggcgatc cggcgtggc ctacggctac gccaggacc aggaaccga cgacgcggc	1380
cggatccgca acggtgccct gctgcgggtc tatgtgcgc gctcgagcct gccgggcttc	1440
taccgcacca gcctgacct ggccgcgcg gaggcggcgg gcgaggtcga acggtgac	1500
ggccatccgc tgccgtcgc cctggacgcc atcaccggcc ccgaggagga agcggggcgc	1560

Sequence Listing

ctggagacca ttctcggtg gccgtggcc gagcgaccg tgggtattcc ctcgcgatc	1620
cccaccgacc cgcgcaacgt cggcggcgac ctcgaccgt ccagcatccc cgacaaggaa	1680
caggcgatca gcgccctgcc ggactacgcc agccagcccg gcaaaccgcc gcgcgaggac	1740
ctgaagtaa	1749

<210> 2

<211> 1764

<212> DNA

<213> Artificial Sequence

<220>

<223> pMH21 plasmid coding sequence

<400> 2

atggaggatga agctgggtga atctggagga ggcttagtgc agcctggagg gtccctgaaa	60
ctctctgtg caacctctg attcactttc agtgactatt acatgtattg ggttcgccag	120
actccagaga agaggctga gtgggtcgca tacattagta atgatgatag ttccgccgct	180
tattcagaca ctgtaaagg cgggttcacc atctccagag acaatgccag gaacaccctc	240
tacctgcaaa tgagccgtct gaagtctgag gacacagcca tatattcctg tgcaagagga	300
ctggcctggg gagcctggtt tgcttactgg ggccaaggga ctctgggtcac tgtctctgca	360
gccaaaacga ccccccatc tgtctatcca ctggccctg gatctgtctc ccaaactaac	420
tccatggtga cctgggatg cctggtcaag ggctatttcc ctgagccagt gacagtgacc	480
tggaactctg gatccctgtc cagcgtgtg cacaccttc cagctgtcct gcagtctgac	540
ctctacactc tgagcagctc agtgactgtc cctccagca cctggcccag cgagaccgtc	600
acctgcaacg ttgccaccc gccagcagc accaagggtg acaagaaaat tgtgccagg	660